

FIGURE 1



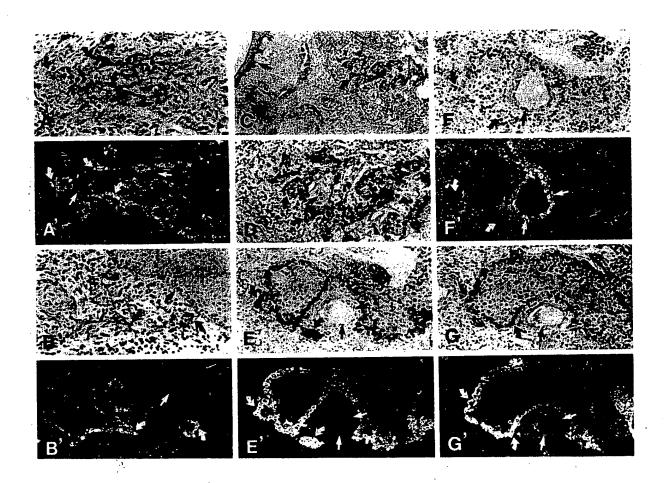


FIGURE 2





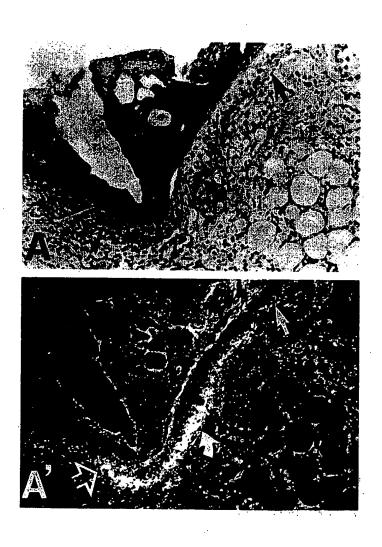


FIGURE 3

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FIG 4A (SEQ ID NO.:12 & 13)

1 gaccacetga tegaaggaaa aggaaggeac ageggagege agagtgagaa ceaceaaceg 61 aggegeeggg cagegaecee tgeageggag acagagaetg ageggeeegg cacegeeatg 1 M
121 cetgegetet ggetgggetg etgeetetge ttetegetee teetgeeege ageeegggee

2 P A L W L G C C L C F S L L L P A A R A

181 acctccagga gggaagtctg tgattgcaat gggaagtcca ggcagtgtat ctttgatcgg
22 T S R R E V C D C N G K S R Q C I F D R

↓⇔ Domain V

241 gaacttcaca gacaaactgg taatggattc cgctgcctca actgcaatga caacactgat 42 E L H R Q T G N G F R C L N C N D N T D 301 ggcattcact gcgagaagtg caagaatggc ttttaccggc acagagaaag ggaccgctgt 62 GIH CEKCKNG FYR HRER DRC 361 ttgccctgca attgtaactc caaaggttct cttagtgctc gatgtgacaa ctctggacgg 82 L P C N C N S K G S L S A R C D N S G R 421 tgcagctgta aaccaggtgt gacaggagec agatgcgacc gatgtctgcc aggettecac 102 C S C K P G V TGARCDRCLP GFH 481 atgeteacgg atgeggggtg cacceaagae cagagaetge tagaetecaa gtgtgaetgt DAGCTQD QRLLDS.K CDC 122 M L T 541 gacccagety geategeagy gecetytyae gegggeeget gtytetyeaa gecagetytt 142 D P A G I A G P C D AGR CVCK 601 actggagaac getgtgatag gtgtcgatca ggttactata atctggatgg ggggaaccet 162 T G E R C D R C R S G Y Y N L D G G N P 661 gagggetgta eccagtgttt etgetatggg catteageca getgeegeag etetgeagaa H S A S C R S S A E 182 E G C T Q C F C Y G !⇒ Domain IV

721 tacagtgtcc ataagatcac ctctaccttt catcaagatg ttgatggctg gaaggctgtc 202 Y S V H K I T S T F H Q D V D G W KAV 781 caacgaaatg ggtctcctgc aaagctccaa tggtcacagc gccatcaaga tgtgtttagc GSPAKLQ WSQRHQD QRN 841 teagecease gactagates tytetatttt gtggeteetg ceasatttet tgggastesa VAP 242 S A Q R L D P VYF AKFL GNO 901 caggtgaget atgggeaaag cetgteettt gactacegtg tggacagagg aggeagacae 262 Q V S Y G Q S L S F D Y R V D R G GRH 961 ccatctgccc atgatgtgat cctggaaggt gctggtctac ggatcacagc tcccttgatg LEGAGL RITA PLM 282 P S A H D V I 1021 ccacttggca agacactgcc ttgtgggctc accaagactt acacattcag gttaaatgag 302 PLG KTLP CGL TKT YTFR LNE 1081 catecaagea ataattggag ecceeagetg agttaetttg agtategaag gttaetgegg S. Y F E Y R R P Q L LLR 322 H P S N N W S 1141 aatotoacag cootoogoat cogagotaca tatggagaat acagtactgg gtacattgac YGE YSTG 342 N L T ALRI RAT 1201 aatgtgacco tgatttcago cogocotgto totggagooo cagoaccotg ggttgaacag 362 N V T L I S A R P V SGAPAPW 1261 tgtatatgtc ctgttgggta caaggggcaa ttctgccagg attgtgcttc tggctacaag DCAS 382 C I C P V G Y KGQ F C Q 1⇒ Domain III

1321 agagattcag cgagactggg gccttttqgc acctgtattc cttgtaactg tcaaggggga 402 R D S A R L G P F G T C I P C N C Q G G 1381 ggggcetgtg atccagacac aggagattgt tattcagggg atgagaatcc tgacattgag 422 G A C D P D T G D C Y S G D E N P D I E 1441 tgtgctgact gcccaattgg tttctacaac gatccgcacg accccgcag ctgcaagcca 442 C A D C P I G F Y N D P H D P R S C K P 1501 tgtccctgtc ataacgggtt cagctgctca gtgattccgg agacggagga ggtggtgtgc 462 C P C H N G F S C S V I P E T E E V V C

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FIG 4B (SEQ ID NO.:12 & 13)

1561 aataactgee eteceggggt caceggtgee egetgtgage tetgtgetga tggetaettt 482 N N C P P G V TGARCE LCAD GY ? 1621 ggggacccct ttggtgaaca tggcccagtg aggccttgtc agccctgtca atgcaacagc G P V RPC QPC Q C N S 502 G D P F G E H 1681 aatgtggacc ccagtgcctc tgggaattgt gaccggctga Caggcaggtg tttgaagtgt GNC DRL TGRC 522 N. V D P S A 5 L K C 1741 atocacaaca cagooggoat otactgogao cagtgoaaag Caggotactt oggggaocca 542 I H N T A G I Y C D Q C K A G Y F G D P 1801 ttggctccca acccagcaga caagtgtcga gcttgcaact gtaaccccat gggctcagag 562 LAPNPAD KCRACN CNPM GSE 1861 cctgtaggat gtcgaagtga tggcacctgt gtttgcaagc caggatttgg tggccccaac 582 PVG CRSD GTC VCK PG FG G PN 1921 tgtgagcatg gagcattcag ctgtccagct tgctataatc aagtgaagat tcagatggat 602 C E H GAFS CPACYN QVKI [→ Domain I/II 1981 cagittatgc agcagettca gagaatggag geeetgattt caaaggetea gggtggtgat 622 Q F M Q Q L Q RME ALI S K A Q G G D 2041 ggagtagtac ctgatacaga gctggaaggc aggatgcagc aggctgagca ggcccttcag LEGRMQ QAEQ ALQ 642 G V V P D T E 2101 gacattetga gagatgeeca gattteagaa ggtgetagea gateeettgg teteeagttg 662 DIL RDAQISE GAS RS LG 2161 gecaaggtga ggagecaaga gaacagetae cagageegee tggatgaeet caagatgaet 682 A K V R S O E N S Y O S R L D D L K M T 2221 qtqqaaaqaq ttcqqqctct qqqaaqtcag taccaqaacc gagttcqqqa tactcacaqq 702 V E R V R A L G S Q Y Q N R V R D T H R 2281 ctcatcactc agatgcagct gagcctggca gaaagtgaag cttccttggg aaacactaac 722 L I T QMQL SLA E S E A S L G NTN 2341 attectgeet cagaceacta egtggggeea aatggettta aaagtetgge teaggaggee 742 I P A S D H Y V G P N G F K S L A Q E A 2401 acaagattag cagaaagcca cgttgagtca gccagtaaca tggagcaact gacaagggaa 762 T R L A E S H VES ASN MEQL TRE 2461 actgaggact attccaaaca agccctctca ctggtgcgca aggccctgca tgaaggagtc 782 T E D Y S  $_{\perp}$  K Q A L S L V R K A L H E G V 782 T E D Y S K Q 2521 ggaageggaa geggtageee ggaeggtget gtggtgeaag ggettgtgga aaaattggag 802 G S G S P D G A V V Q G L V E K L E 2581 aaaaccaagt ccctggccca gcagttgaca agggaggcca ctcaagcgga aattgaagca 822 KTK SLAQ Q L'T REATQAE I E A 2641 gataggicti atcagcacag totocquoto otqqattoag igiotocqui toagggagto VSPL 842 D R S Y Q H S LRL L D S 0 G 2701 agtgatcagt cctttcaggt ggaagaagca aagaggatca aacaaaaagc ggattcactc SDQSFQV EEA KRI KQKA DSL 2761 tcaagcctgg taaccaggca tatggatgag ttcaagcgta cacaaaagaa tctgggaaac 882 S S L V T R H M D E F K R T Q K N L G N 2821 tggaaagaag aagcacagca gotottacag aatggasaaa gtgggagaga gaaatcagat LLQ NGK SGRE KSD 902 W K E E A Q Q 2881 cagetgettt ecegtgecaa tettgetaaa ageagageae aagaageaet gagtatggge 922 Q L L S R A N L A K S R A Q E A L S M G 2941 aatgccactt tttatgaagt tgagagcatc cttaaaaacc tcagagagtt tgacctgcag 942 NAT FYEVESI LKN LREF D L Q 3001 gtggacaaca gaaaagcaga agctgaagaa gccatgaaga gactctccta catcagccag 962 V D N R K A E A E E A M K R L S Y I S Q 3061 aaggtttcag atgccagtga caagacccag caagcagaaa gagccctggg gagcgctgct 982 K V S D A S D K T Q Q A E R A L G S A A 3121 gctgatgcac agagggcaaa gaatggggcc ggggaggccc tggaaatctc cagtgagatt Q R A K N G A G E A L E I S S E I 1002 A D A



FIG 4C (SEQ ID NO.: 12 & 13) 3181 gaacaggaga ttgggagtct gaacttggaa gccaatgtga cagcagatgg agccttggcc IGSL N L E A N V T A D G A L A 3241 atggaaaagg gactggcctc tctgaagagt gagatgaggg aagtggaagg agagctggaa EVEG LKS E M R GLAS 1042 M E K 3301 aggaaggage tggagtttga cacqaatatg gatgcagtac agatggtgat tacagaagce DAV QM·VI 1062 R K E L E F D T N M 3361 cagaaggttg ataccagage caagaaeget ggggttacaa tecaagaeae acteaaeaea GVTIQDT 1082 Q K V D T R A K N A 3421 tragacggcc tectgeater gatggaccag ceteteagtg tagatgaaga ggggetggte M D Q PLSVDEE GLV 1102 L D G L L H L 3481 ttactggage agaagettte cegagecaag acceagatea acagecaact geggeceatg RAKTQINSQL 1122 L L E Q K L S 3541 atgtcagago tggaagagag ggcacgtcag cagaggggcc acctccattt gctggagaca 1142 M S E L E E R A R Q Q R G H L H L 3601 agcatagatg ggattetgge tgatgtgaag aacttggaga acattaggga caacetgeee NLENIRD NLP GILADVK 1162 S I D 3661 ccaggetget acaataccca ggetettgag caacagtgaa getgecataa atatttetca ALE QQ YNTQ 3721 actgaggttc ttgggataca gatctcaggg ctcgggagcc atgtcatgtg agtgggtggg 3781 atggggacat ttgaacatgt ttaatgggta tgctcaggtc aactgacctg accccattcc 3841 tgatcccatg gccaggtggt tgtcttattg caccatactc cttgcttcct gatgctgggc 3901 atgaggcaga taggcactgg tgtgagaatg atcaaggatc tggaccccaa agatagactg 3961 gatggaaaga caaactgcac aggcagatgt ttgcctcata atagtcgtaa gtggagtcct 4021 ggaatttgga caagtgetgt tgggatatag teaacttatt etttgägtaa tgtgaetaaa 4081 ggaaaaact ttgactttgc ccaggcatga aattcttcct aatgtcagaa cagagtgcaa 4141 cocagtoaca ctgtggccag taaaatacta ttgcctcata ttgtcctctg caagettctt 4201 getgateaga gtteeteeta ettacaacce agggtgtgaa catgttetee atttteaage 4261 tggaagaagt gagcagtgtt ggagtgagga cotgtaaggc aggcccattc agagctatgg 4321 tgcttgctgg tgcctgccac cttcaagttc tggacctggg catgacatcc tttcttttaa 4441 atgttgggaa agtatttact ttttcggttt caaagtgata gaaaagtgtg gcttgggcat 4501 tgaaagaggt aasattetet agatttatta gteetaatte aateetaett ttegaacaee 4561 aaaaatgatg cgcatcaatg tattttatct tattttctca atctcctctc tctttcctcc 4621 acccataata agagaatgtt cctactcaca cttcagctgg gtcacatcca tccctccatt 4681 catectteca tecatette catecattae etecatecat cettecaaca tatatttatt 4741 gagtacctac tgtgtgccag gggctggtgg gacagtggtg acatagtctc tgccctcata 4801 gagttgattg tctagtgagg aagacaagca tttttaaaaa ataaatttaa acttacaaac 4861 tetgetegec acaageggeg tetategcaa taaccgceeg geetgcaacc tetetgceca 4921 acagaacata tgttgcaaga cootcocatg ggcactgagt ttggcaagga tgacagagct 4981 ctgggttgtg cacatttett tgeattecag cgteactetg tgeettetae aactgattge 5041 aacagactgt tgagttatga taacaccagt gggaattget ggaggaacca gaggcactte 5101 caccttggct gggaagacta tggtgctqcc ttgcttctgt atttccttgg attttcctga

5161 aagtgttttt aaataaagaa caattgttag atgccaaaaa //



FIG 4D (SEQ ID NO.:14 & 15) 3421 ttagacggcc tectgeatet gatgggtatg tgaacceaca acceacaacc ttecagetee 1102 L D G L L H L M G M 3481 atgetecagg getttgetee agaacaetea ctatacetag ceccageaaa ggggagtete 3541 agettteett aaggatatea gtaaatgtge titgtiteea ggeecagata acttreggea 3601 ggttccctta catttactgg accetgtttt accgttgcta agatgggtca ctgaacacet 3661 attgcacttg ggggtaaagg totgtgggcc aaagaacagg tgtatataag caacttcaca 3721 gaacacgaga cagettggga atcetgetaa agagtetgge etggaccetg agaagecagt 3781 ggacagtttt aagcagagga ataacatcac cacigtatat ttcagaaaga tcactagggc 3841 agccgagtgg aggaaagctt gaagaggggg ttagagagaa ggcaggttga gactacttaa 3901 gatattgttg aaataattga agagagaaat gacaggagcc tgctctaagg cagtagaatg 3961 gtggctggga agatgtgaag gaagattttc ccagtctgtg aagtcaagaa tcacttgccq 4021 gccgggtgtg gtggctcacg cctgtaattc tagcactttg ggagactgaa gcgggtggat 4081 caccogaggt caggagttga agaccagcot ggccaacatg gtgaaaccot gtototacta 4141 aaagtacaaa aattagotgg atgatggtgg tgggcgcotg taattocago tactcaggag 4201 totgaggoag gagaatogot tgaaccoagg aggogaggot acagtgagoo aagattgoac 4261 cactgotott coagootggg aacagagaga otgootaaaa aaaaaaaaaa aaaaaa //



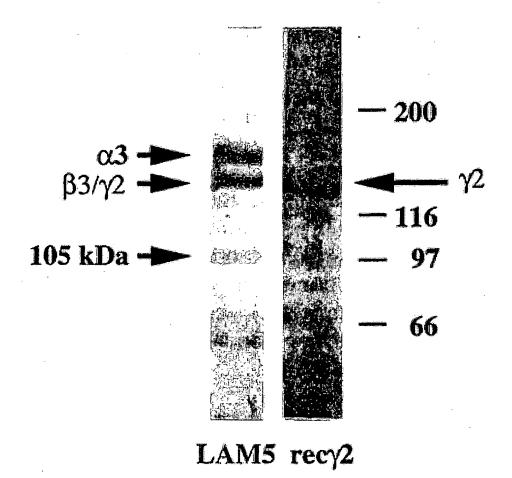


FIGURE 5



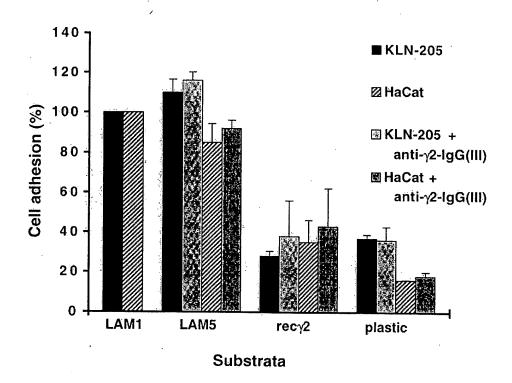
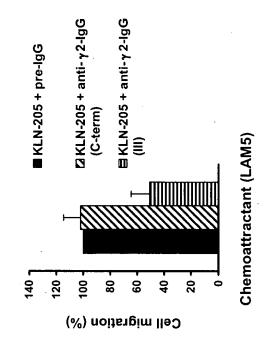
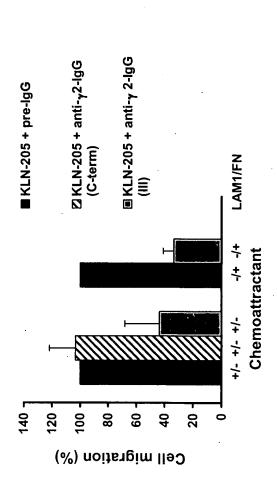


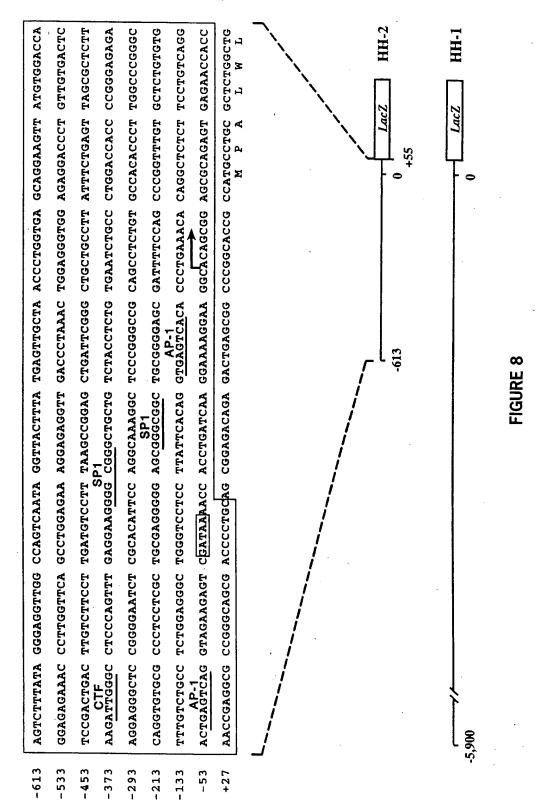
FIGURE 6













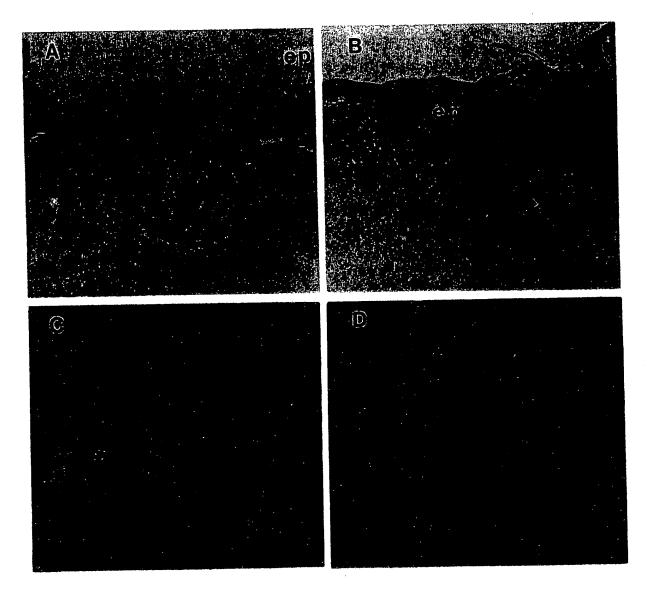


FIGURE 11



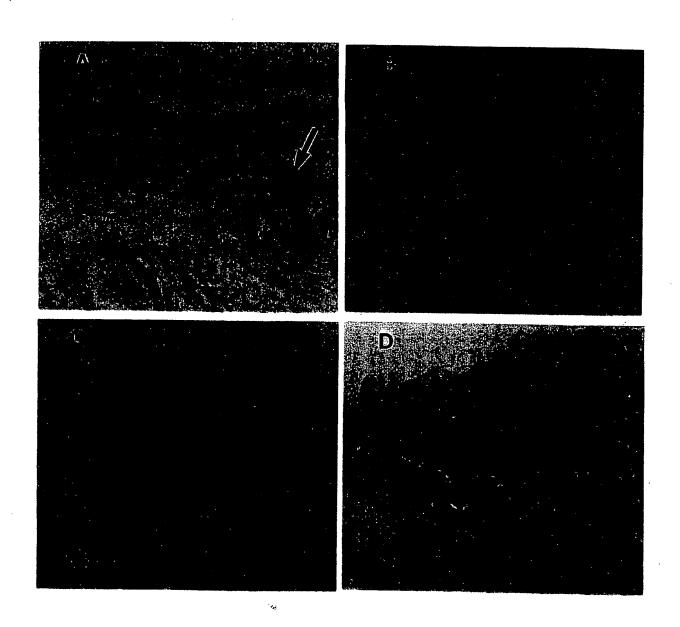


FIGURE 10



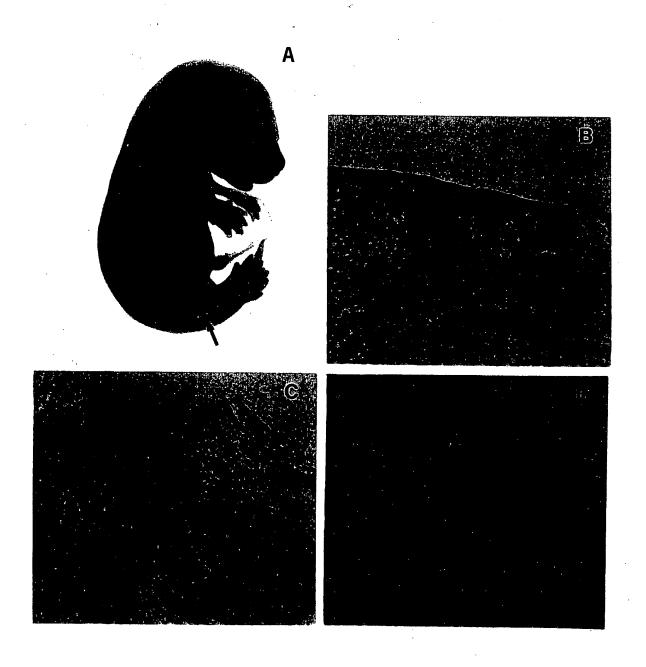


FIGURE 9